

HSU44805

LOCUS HSU44805 1979 bp DNA linear PRI 02-FEB-1996

DEFINITION Human prostaglandin H synthase type 2 (PHS-2) gene, promoter sequence and partial cds.

ACCESSION U44805

VERSION U44805.1 GI:1174223

KEYWORDS

SOURCE Homo sapiens (human)

ORGANISM Homo sapiens

Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.

REFERENCE 1 (bases 1 to 1979)

AUTHORS Kutchera, W.A., Jones, D.A., Matsunami, N., Groden, J., McIntyre, T.M.,
 Zimmerman, G.A., White, R.L. and Prescott, S.M.

TITLE Prostaglandin H synthase-2 is expressed abnormally in human colon cancer: evidence for a transcriptional effect

JOURNAL Proc. Natl. Acad. Sci. U.S.A. (1996) In press

REFERENCE 2 (bases 1 to 1979)

AUTHORS Kutchera, W.A.

TITLE Direct Submission

JOURNAL Submitted (05-JAN-1996) William A. Kutchera, Human Molecular
Biology and Genetics, University of Utah, Building 533, Salt Lake
City, UT 84112, USA

FEATURES

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source      1. .1979
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            /db_xref="taxon:9606"
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gene      1. .1979
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promoter 1. .1976
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CDS      1977. .>1979
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          /protein_id="AAD14842.1"
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ORIGIN

Query Match 97.3%; Score 1851; DB 9; Length 1979;

Best Local Similarity 99.9%; Pred. No. 0;

Matches 1901; Conservative 0; Mismatches 1; Indels 0; Gaps 0;

Qy 1 GGATTCTAACATGGCTTCTAACCCAAACTAACATTAGTAGCTCTAACTATAAACTTCAA 60
| | | | |

Db 46 GGATTCTAACATGGCTTCTAACCCAAACTAACATTAGTAGCTCTAACTATAAACTTCAAA 105

Qy 61 TTTCAGTAGATGCAACCTACTCCTTTAAATGAAACAGAAGATTGAAATTATTAAATTAT 120
 | | | | | | | | | | | | | | | | | | | | | | | | | | | |

Db 106 TTT CAG TAG ATG CAAC CTA CTC CTT TAAA ATG AAAC AGA AGATTG AAATT ATTA AAATTAT 165

| | | | |
|----|-----|---|------|
| Qy | 121 | CAAAAAAGAAAATGATCCACGCTCTTAGTTGAAATTTTCATGTAAGATTCCATGCAATAAAAT | 180 |
| Db | 166 | CAAAAAAGAAAATGATCCACGCTCTTAGTTGAAATTTTCATGTAAGATTCCATGCAATAAAAT | 225 |
| Qy | 181 | AGGAGTGCCATAAAATGGAATGATGAAATATGACTAGAGGAGGAGAAAGGCTTCCTAGATG | 240 |
| Db | 226 | AGGAGTGCCATAAAATGGAATGATGAAATATGACTAGAGGAGGAGAAAGGCTTCCTAGATG | 285 |
| Qy | 241 | AGATGGAATTTTGTATCATCCGTGTCTCATGAAGAATCAGATGTGTACACTAAGCAAAAACA | 300 |
| Db | 286 | AGATGGAATTTTGTATCATCCGTGTCTCATGAAGAATCAGATGTGTACACTAAGCAAAAACA | 345 |
| Qy | 301 | GTTAAAAAAGAAACCTCCAAGTGAGTCTCTTATTTATTTTTTCTTATAAGACTTCTACA | 360 |
| Db | 346 | GTTAAAAAAGAAACCTCCAAGTGAGTCTCTTATTTATTTTTTCTTATAAGACTTCTACA | 405 |
| Qy | 361 | AATTGAGGTACCTGGTGTAGTTTTATTTTCAGGTTTTATGCTGTCATTTTCCTGTAATGCT | 420 |
| Db | 406 | AATTGAGGTACCTGGTGTAGTTTTATTTTCAGGTTTTATGCTGTCATTTTCCTGTAATGCT | 465 |
| Qy | 421 | AAGGACTTAGGACATAACTGAATTTTCTATTTTCCACTTCTTTTCTGGTGTGTGTGTATA | 480 |
| Db | 466 | AAGGACTTAGGACATAACTGAATTTTCTATTTTCCACTTCTTTTCTGGTGTGTGTGTATA | 525 |
| Qy | 481 | TATATATGTATATATACACACACACATATACATATATATATTTTTTAGTATCTCACCTCA | 540 |
| Db | 526 | TATATATGTATATATATACACACACACATATACATATATATATTTTTTAGTATCTCACCTCA | 585 |
| Qy | 541 | CATGCTCCTCCCTGAGCACTACCCATGATAGATGTTAAACAAAAGCAAAGATGAAATTCC | 600 |
| Db | 586 | CATGCTCCTCCCTGAGCACTACCCATGATAGATGTTAAACAAAAGCAAAGATGAAATTCC | 645 |
| Qy | 601 | AACTGTTAAAATCTCCCTTCCATCTAATTAATTCCTCATCCAACATGTTCCAAAACGAG | 660 |
| Db | 646 | AACTGTTAAAATCTCCCTTCCATCTAATTAATTCCTCATCCAACATGTTCCAAAACGAG | 705 |
| Qy | 661 | AATAGAAAATTAGCCCCAATAAGCCCAGGCAACTGAAAAGTAAATGCTATGTTGTACTTT | 720 |
| Db | 706 | AATAGAAAATTAGCCCCAATAAGCCCAGGCAACTGAAAAGTAAATGCTATGTTGTACTTT | 765 |
| Qy | 721 | GATCCATGGTCACAACCTCATAATCTTGGAAGAGTGGACAGAAAAGACAAAAGAGTGAAC | 780 |
| Db | 766 | GATCCATGGTCACAACCTCATAATCTTGGAAGAGTGGACAGAAAAGACAAAAGAGTGAAC | 825 |
| Qy | 781 | TTAAAACCTCGAATTTATTTTACCAGTATCTCCTATGAAGGGCTAGTAACCAAAATAATCC | 840 |
| Db | 826 | TTAAAACCTCGAATTTATTTTACCAGTATCTCCTATGAAGGGCTAGTAACCAAAATAATCC | 885 |
| Qy | 841 | ACGCATCAGGGAGAGAAATGCCTTAAGGCATACGTTTGGACATTTAGCGTCCCTGCAAA | 900 |
| Db | 886 | ACGCATCAGGGAGAGAAATGCCTTAAGGCATACGTTTGGACATTTAGCGTCCCTGCAAA | 945 |
| Qy | 901 | TTCTGGCCATCGCCGCTTCCTTTGTCCATCAGAAGGCAGGAACTTTATATTGGTGACCC | 960 |
| Db | 946 | TTCTGGCCATCGCCGCTTCCTTTGTCCATCAGAAGGCAGGAACTTTATATTGGTGACCC | 1005 |
| Qy | 961 | GTGGAGCTCACATTAACCTATTTTACAGGGTAACTGCTTAGGACCAGTATTATGAGGAGGAT | 1020 |

| | | | |
|----|------|--|------|
| Db | 1006 | GTGGAGCTCACATTAACCTATTTACAGGGTAACTGCTTAGGACCAGTATTATGAGGAGAAT | 1065 |
| Qy | 1021 | TTACCTTTCCCGCCTCTCTTTTCCAAGAAACAAGGAGGGGGTGAAGGTACGGAGAACAGTA | 1080 |
| Db | 1066 | TTACCTTTCCCGCCTCTCTTTTCCAAGAAACAAGGAGGGGGTGAAGGTACGGAGAACAGTA | 1125 |
| Qy | 1081 | TTTCTTCTGTTGAAAGCAACTTAGCTACAAAGATAAATTACAGCTATGTACACTGAAGGT | 1140 |
| Db | 1126 | TTTCTTCTGTTGAAAGCAACTTAGCTACAAAGATAAATTACAGCTATGTACACTGAAGGT | 1185 |
| Qy | 1141 | AGCTATTTTCATTCCACAAAATAAGAGTTTTTTTAAAAAGCTATGTATGTATGTCCTGCATA | 1200 |
| Db | 1186 | AGCTATTTTCATTCCACAAAATAAGAGTTTTTTTAAAAAGCTATGTATGTATGTCCTGCATA | 1245 |
| Qy | 1201 | TAGAGCAGATATACAGCCTATTAAGCGTCGTCACTAAACATAAAACATGTCAGCCTTTC | 1260 |
| Db | 1246 | TAGAGCAGATATACAGCCTATTAAGCGTCGTCACTAAACATAAAACATGTCAGCCTTTC | 1305 |
| Qy | 1261 | TTAACCTTACTCGCCCCAGTCTGTCCCGACGTGACTTCCTCGACCCTCTAAAGACGTACA | 1320 |
| Db | 1306 | TTAACCTTACTCGCCCCAGTCTGTCCCGACGTGACTTCCTCGACCCTCTAAAGACGTACA | 1365 |
| Qy | 1321 | GACCAGACACGGCGGGCGGGCGGGAGAGGGGATTCCCTGCGCCCCCGGACCTCAGGGCC | 1380 |
| Db | 1366 | GACCAGACACGGCGGGCGGGCGGGAGAGGGGATTCCCTGCGCCCCCGGACCTCAGGGCC | 1425 |
| Qy | 1381 | GCTCAGATTCTGAGAGGAAGCCAAGTGTCTTCTGCCCTCCCCCGGTATCCCATCCAA | 1440 |
| Db | 1426 | GCTCAGATTCTGAGAGGAAGCCAAGTGTCTTCTGCCCTCCCCCGGTATCCCATCCAA | 1485 |
| Qy | 1441 | GGCGATCAGTCCAGAACTGGCTCTCGGAAGCGCTCGGGCAAAGACTGCGAAGAAGAAAAG | 1500 |
| Db | 1486 | GGCGATCAGTCCAGAACTGGCTCTCGGAAGCGCTCGGGCAAAGACTGCGAAGAAGAAAAG | 1545 |
| Qy | 1501 | ACATCTGGCGGAAACCTGTGCGCCTGGGGCGGTGGAACCTCGGGGAGGAGAGGGAGGGATC | 1560 |
| Db | 1546 | ACATCTGGCGGAAACCTGTGCGCCTGGGGCGGTGGAACCTCGGGGAGGAGAGGGAGGGATC | 1605 |
| Qy | 1561 | AGACAGGAGAGTGGGGACTACCCCTCTGCTCCCAAATTGGGGCAGCTTCTGGGTTTCC | 1620 |
| Db | 1606 | AGACAGGAGAGTGGGGACTACCCCTCTGCTCCCAAATTGGGGCAGCTTCTGGGTTTCC | 1665 |
| Qy | 1621 | GATTTTCTCATTTCCGTGGGTAAAAAACCTGCCCCACCGGGCTTACGCAATTTTTTTTA | 1680 |
| Db | 1666 | GATTTTCTCATTTCCGTGGGTAAAAAACCTGCCCCACCGGGCTTACGCAATTTTTTTTA | 1725 |
| Qy | 1681 | AGGGGAGAGGAGGGGAAAAATTTGTGGGGGGTACGAAAAGGCGGAAAGAAACAGTCATTTT | 1740 |
| Db | 1726 | AGGGGAGAGGAGGGGAAAAATTTGTGGGGGGTACGAAAAGGCGGAAAGAAACAGTCATTTT | 1785 |
| Qy | 1741 | GTCACATGGGCTTGGTTTTTCAGTCTTATAAAAAGGAAGGTTCTCTCGGTTAGCGACCAAT | 1800 |
| Db | 1786 | GTCACATGGGCTTGGTTTTTCAGTCTTATAAAAAGGAAGGTTCTCTCGGTTAGCGACCAAT | 1845 |
| Qy | 1801 | TGTCATACGACTTGACAGTGAGCGTCAGGAGCACGTCCAGGAACCTCCTCAGCAGCGCCTCC | 1860 |

Db 1846 TGTACATACGACTTGTCAGTGAGCGTCAGGAGCACGTCCAGGAACCTCCTCAGCAGCGCCTCC 1905

Qy 1861 TTCAGCTCCACAGCCAGACGCCCTCAGACAGCAAAGCCTACC 1902

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Db 1906 TTCAGCTCCACAGCCAGACGCCCTCAGACAGCAAAGCCTACC 1947